

Amendments to the Claims:

This listing of claims will replace all prior versions, and listings of claims in the application:

LISTING OF CLAIMS:

1. (Original) A method of forming a peptide conjugate comprising a covalent linkage between a modifying group and a glycosylated or non-glycosylated peptide, wherein said modifying group is conjugated to the peptide via a glycosyl linking group interposed between and covalently linked to both said peptide and said modifying group, said method comprising:
 - a. contacting a cell with a modified sugar comprising a sugar moiety and at least one modifying group, wherein said modifying group is a member independently selected from the group consisting of a water-soluble polymer, a therapeutic moiety, a detectable label, a biomolecule and a targeting moiety;
 - b. incubating said cell under conditions in which said cell internalizes said modified sugar;
 - c. after step b, intracellularly contacting said modified sugar with a glycosylated or non-glycosylated peptide and a glycosyltransferase for which said modified sugar is a substrate, thereby forming said peptide conjugate.
2. (Original) The method of claim 1, further comprising, after step b and before step c, intracellularly contacting said modified sugar with a nucleotide and a nucleotidyl transferase, thereby forming a modified nucleotide sugar, wherein said modified sugar in step c is said modified nucleotide sugar.
3. (Original) The method of claim 1, further comprising isolating said peptide conjugate.
4. (Original) The method of claim 1, wherein said modified sugar is a modified nucleotide sugar.
5. (Original) The method of claim 1, wherein said modified sugar is a modified activated sugar.

6. (Original) The method of claim 1, wherein said glycosyl linking group is an intact glycosyl linking group.

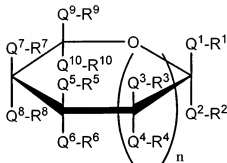
7. (Original) The method of claim 1, wherein said modified sugar is a precursor modified sugar that is intracellularly converted to an intermediate modified sugar by cellular enzymes after step b and before step c.

8. (Original) The method of claim 7, wherein said intermediate modified sugar is a phosphorylated modified sugar, wherein said phosphorylated modified sugar is formed by intracellularly contacting said modified sugar with a kinase for which said modified sugar is a substrate, thereby forming a phosphorylated modified nucleotide sugar.

9. (Original) The method of claim 1, wherein said water-soluble polymer comprises poly(ethylene glycol).

10. (Original) The method of claim 10, wherein said poly(ethylene glycol) has a molecular weight distribution that is essentially homodisperse.

11. (Original) The method of claim 1, wherein said modified sugar has the formula



(I)

wherein,

n represents an integer from 0 to 1;

Q^1 , Q^2 , Q^3 , Q^4 , Q^5 , Q^6 , Q^7 , Q^8 , Q^9 , and Q^{10} are members independently selected from a bond, substituted or unsubstituted alkylene, substituted or unsubstituted heteroalkylene, substituted or unsubstituted cycloalkylene, substituted or unsubstituted heterocycloalkylene, substituted or unsubstituted

arylene, substituted or unsubstituted heteroarylene, -O-, -N(R^{1A})-, -S-, -C(O)-, and -CH₂-, wherein
R^{1A} is a member selected from hydrogen, substituted or unsubstituted alkyl, substituted or unsubstituted heteroalkyl, substituted or unsubstituted cycloalkyl, substituted or unsubstituted heterocycloalkyl, substituted or unsubstituted aryl, and substituted or unsubstituted heteroaryl; and
R¹, R², R³, R⁴, R⁵, R⁶, R⁷, R⁸, R⁹, and R¹⁰ are members independently selected from -OPO₃H₂, -OH, -NH₂, -SH, hydrogen, substituted or unsubstituted alkyl, substituted or unsubstituted heteroalkyl, substituted or unsubstituted cycloalkyl, substituted or unsubstituted heterocycloalkyl, substituted or unsubstituted aryl, substituted or unsubstituted heteroaryl, an activated leaving group, a nucleotidyl moiety, and a modifying group, wherein at least one of R¹, R², R³, R⁴, R⁵, R⁶, R⁷, R⁸, R⁹, and R¹⁰ is a modifying group.

12. (Original) The method of claim 11, wherein

Q¹-R¹, Q²-R², Q³-R³, Q⁴-R⁴, Q⁵-R⁵, Q⁶-R⁶, Q⁷-R⁷, Q⁸-R⁸, Q⁹-R⁹, and Q¹⁰-R¹⁰ are members independently selected from hydrogen, -OPO₃H₂, -OH, -OCH₃, -CH₃, -C(O)H, -CH₂OH, -NHR¹¹, -O-CH(CH₃)COOR¹², -C(O)OR¹³, -CHR¹⁴-CHR¹⁵-CH₂R¹⁶, an activated leaving group, a nucleotidyl moiety and -L-M, wherein at least one of R¹, R², R³, R⁴, R⁵, R⁶, R⁷, R⁸, R⁹, and R¹⁰ is -L-M, wherein

L is a linker independently selected from a bond, substituted or unsubstituted alkylene, substituted or unsubstituted heteroalkylene, substituted or unsubstituted cycloalkylene, substituted or unsubstituted heterocycloalkylene, substituted or unsubstituted arylene, substituted or unsubstituted heteroarylene, -O-, -NH-, -S-, and CH₂-,

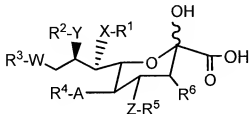
M is a modifying group, and

R¹¹, R¹², R¹³, R¹⁴, R¹⁵, and R¹⁶ are independently selected from hydrogen, substituted or unsubstituted alkyl, substituted or unsubstituted heteroalkyl, and -L¹-M¹, wherein

L¹ is a linker independently selected from a bond, substituted or unsubstituted alkylene, substituted or unsubstituted heteroalkylene,

substituted or unsubstituted cycloalkylene, substituted or unsubstituted heterocycloalkylene, substituted or unsubstituted arylene, substituted or unsubstituted heteroarylene, -O-, -NH-, -S-, and CH₂-, and M¹ is modifying group.

13. (Original) The method of claim 11, wherein said modified sugar has the formula

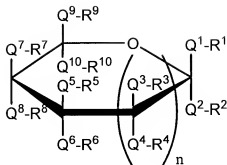


(II)

wherein,

W, X, Y, Z, and A are members independently selected from a bond, substituted or unsubstituted alkylene, substituted or unsubstituted heteroalkylene, substituted or unsubstituted cycloalkylene, substituted or unsubstituted heterocycloalkylene, substituted or unsubstituted arylene, substituted or unsubstituted heteroarylene, -O-, -N(R⁷), -S-, and -CH₂-, wherein, R⁷ is a member independently selected from hydrogen, substituted or unsubstituted alkyl, substituted or unsubstituted heteroalkyl, substituted or unsubstituted cycloalkyl, substituted or unsubstituted heterocycloalkyl, substituted or unsubstituted aryl, and substituted or unsubstituted heteroaryl; and R¹, R², R³, R⁴, R⁵ and R⁶ are members independently selected from -OH, -NH₂, -SH, hydrogen, substituted or unsubstituted alkyl, substituted or unsubstituted heteroalkyl, substituted or unsubstituted cycloalkyl, substituted or unsubstituted heterocycloalkyl, substituted or unsubstituted aryl, substituted or unsubstituted heteroaryl, and a modifying group, wherein at least one or R¹, R², R³, R⁴, R⁵ and R⁶ is a modifying group.

14. (Original) The method of claim 4, wherein said modified nucleotide sugar has the formula



(1)

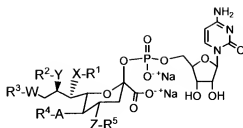
wherein,

n represents an integer from 0 to 1;

Q¹, Q², Q³, Q⁴, Q⁵, Q⁶, Q⁷, Q⁸, Q⁹, and Q¹⁰ are members independently selected from a bond, substituted or unsubstituted alkylene, substituted or unsubstituted heteroalkylene, substituted or unsubstituted cycloalkylene, substituted or unsubstituted heterocycloalkylene, substituted or unsubstituted arylene, substituted or unsubstituted heteroarylene, -O-, -N(R^{1A})-, -S-, -C(O)-, and -CH₂-, wherein

R^{1A} is a member selected from hydrogen, substituted or unsubstituted alkyl, substituted or unsubstituted heteroalkyl, substituted or unsubstituted cycloalkyl, substituted or unsubstituted heterocycloalkyl, substituted or unsubstituted aryl, and substituted or unsubstituted heteroaryl; and R¹, R², R³, R⁴, R⁵, R⁶, R⁷, R⁸, R⁹, and R¹⁰ are members independently selected from -OPO₃H₂, -OH, -NH₂, -SH, hydrogen, substituted or unsubstituted alkyl, substituted or unsubstituted heteroalkyl, substituted or unsubstituted cycloalkyl, substituted or unsubstituted heterocycloalkyl, substituted or unsubstituted aryl, substituted or unsubstituted heteroaryl, an activated leaving group, a nucleotidyl moiety, and a modifying group, wherein at least one of R¹, R², R³, R⁴, R⁵, R⁶, R⁷, R⁸, R⁹, and R¹⁰ is a modifying group and a nucleotidyl moiety.

15. (Original) The method of claim 14, wherein said modified nucleotide sugar has the formula



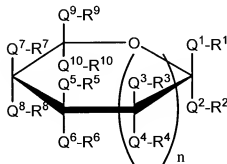
(III)

wherein,

W, X, Y, Z, and A are members independently selected from a bond, substituted or unsubstituted alkylene, substituted or unsubstituted heteroalkylene, substituted or unsubstituted cycloalkylene, substituted or unsubstituted heterocycloalkylene, substituted or unsubstituted arylene, substituted or unsubstituted heteroarylene, -O-, -N(R⁷)-, -S-, and -CH₂-, wherein, R⁷ is a member independently selected from hydrogen, substituted or unsubstituted alkyl, substituted or unsubstituted heteroalkyl, substituted or unsubstituted cycloalkyl, substituted or unsubstituted heterocycloalkyl, substituted or unsubstituted aryl, and substituted or unsubstituted heteroaryl; and

R¹, R², R³, R⁴, and R⁵ are independently selected from -OH, -NH₂, -SH, hydrogen, substituted or unsubstituted alkyl, substituted or unsubstituted heteroalkyl, substituted or unsubstituted cycloalkyl, substituted or unsubstituted heterocycloalkyl, substituted or unsubstituted aryl, substituted or unsubstituted heteroaryl, and a modifying group, wherein at least one of R¹, R², R³, R⁴, and R⁵ is a modifying group.

16. (Original) The method of claim 5, wherein said modified nucleotide sugar has the formula



(I)

wherein,

n represents an integer from 0 to 1;

$Q^1, Q^2, Q^3, Q^4, Q^5, Q^6, Q^7, Q^8, Q^9$, and Q^{10} are members independently selected from a bond, substituted or unsubstituted alkylene, substituted or unsubstituted heteroalkylene, substituted or unsubstituted cycloalkylene, substituted or unsubstituted heterocycloalkylene, substituted or unsubstituted arylene, substituted or unsubstituted heteroarylene, -O-, -N(R^{1A})-, -S-, -C(O)-, and -CH₂-, wherein

R^{1A} is a member selected from hydrogen, substituted or unsubstituted alkyl, substituted or unsubstituted heteroalkyl, substituted or unsubstituted cycloalkyl, substituted or unsubstituted heterocycloalkyl, substituted or unsubstituted aryl, and substituted or unsubstituted heteroaryl; and

R¹, R², R³, R⁴, R⁵, R⁶, R⁷, R⁸, R⁹, and R¹⁰ are members independently selected from -OPO₃H₂, -OH, -NH₂, -SH, hydrogen, substituted or unsubstituted alkyl, substituted or unsubstituted heteroalkyl, substituted or unsubstituted cycloalkyl, substituted or unsubstituted heterocycloalkyl, substituted or unsubstituted aryl, substituted or unsubstituted heteroaryl, an activated leaving group, a nucleotidyl moiety, and a modifying group, wherein at least one of R¹, R², R³, R⁴, R⁵, R⁶, R⁷, R⁸, R⁹, and R¹⁰ is a modifying group and an activated leaving group.

17. (Original) The method of claim 1, wherein said peptide is selected from the group consisting of granulocyte colony stimulating factor, interferon-alpha, interferon-beta, Factor VIIa, Factor IX, follicle stimulating hormone, erythropoietin, granulocyte macrophage colony stimulating factor, interferon-gamma, alpha-1-protease inhibitor, glucocerebrosidase, tissue plasminogen activator protein, interleukin-2, Factor VIII, chimeric tumor necrosis factor receptor, urokinase, chimeric anti-glycoprotein IIb/IIIa antibody, chimeric anti-HER2 antibody, chimeric anti-respiratory syncytial virus antibody, chimeric anti-CD20 antibody, DNase, chimeric anti-tumor necrosis factor antibody, human insulin, hepatitis B sAg, interferon-omega, alpha-galactosidase A, alpha-iduronidase, anti-thrombin III, human chorionic gonadotropin, and human growth hormone.

- 1 **18.** (Original) A cell comprising a peptide conjugate, said peptide conjugate comprising:
2 (i) a modifying group and a peptide, wherein said modifying group is linked to said
3 peptide via a glycosyl linking group interposed between and covalently linked to
4 both the peptide and said modifying group; and
5 (ii) said modifying group is a member independently selected from the group consisting
6 of a water-soluble polymer, a therapeutic moiety, a detectable label, and a
7 targeting moiety.
- 1 **19.** (Original) The method of claim **18**, wherein said glycosyl linking group is an intact
2 glycosyl linking group.